

DATE: 10/07/2002

# 10



OIPE

```
PATENT APPLICATION: US/10/053,520
                                                             TIME: 11:46:22
                    Input Set : A:\46004seq.txt
                    Output Set: N:\CRF4\10072002\J053520.raw
     3 <110> APPLICANT: Sloan-Kettering Institute for Cancer Research
             Rothman, James E.
     5
             Mayhew, Mark
             Hoe, Mee H.
     6 .
     7
             Houghton, Alan
             Hartl, Ulrich
     8
     9
             Ouerfelli, Ouathek
            Moroi, Yoichi
    12 <120> TITLE OF INVENTION: CONJUGATE HEAT SHOCK PROTEIN-BINDING PEPTIDES
    14 <130> FILE REFERENCE: 11746/46004
    16 <140> CURRENT APPLICATION NUMBER: 10/053,520
C--> 17 <141> CURRENT FILING DATE: 2002-10-01
                                                               ENTERED
    19 <150> PRIOR APPLICATION NUMBER: 08/961,707
    20 <151> PRIOR FILING DATE: 1997-10-31
    22 <160> NUMBER OF SEQ ID NOS: 321
    24 <170> SOFTWARE: WordPerfect 8.0 for Windows
    26 <210> SEQ ID NO: 1
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    28 <212> TYPE: PRT
    29 <213> ORGANISM: Artificial Sequence
    31 <220> FEATURE:
    32 <223> OTHER INFORMATION: peptide in m13 coliphage
    34 <400> SEQUENCE: 1
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    37
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    39 <210> SEQ ID NO: 2
    40 <211> LENGTH: 8
    41 <212> TYPE: PRT
    42 <213> ORGANISM: Artificial Sequence
    44 <220> FEATURE:
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    47 <400> SEQUENCE: 2
    49
         Thr Glu Thr Pro Tyr Pro Thr Gly
    50
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    52 <210> SEQ ID NO: 3
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    54 <212> TYPE: PRT
    55 <213> ORGANISM: Artificial Sequence
    57 <220> FEATURE:
    58 <223> OTHER INFORMATION: peptide in m13 coliphage
    60 <400> SEQUENCE: 3
    62
         Leu Thr Thr Pro Phe Ser Ser Gly
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RAW SEQUENCE LISTING

63

Input Set : A:\46004seq.txt

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65 <210> SEQ ID NO: 4
66 <211> LENGTH: 8
67 <212> TYPE: PRT
68 <213> ORGANISM: Artificial Sequence
70 <220> FEATURE:
71 <223> OTHER INFORMATION: peptide in m13 coliphage
73 <400> SEQUENCE: 4
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76
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78 <210> SEQ ID NO: 5
79 <211> LENGTH: 8
80 <212> TYPE: PRT
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: peptide in m13 coliphage
86 <400> SEQUENCE: 5
88 Lys Leu Pro Thr Val Leu Arg Gly
89
      1
91 <210> SEQ ID NO: 6
92 <211> LENGTH: 8
93 <212> TYPE: PRT
94 <213> ORGANISM: Artificial Sequence
96 <220> FEATURE:
97 <223> OTHER INFORMATION: peptide in m13 coliphage
99 <400> SEQUENCE: 6
101 Cys Arg Phe His Gly Asn Arg Gly
      1
104 <210> SEQ ID NO: 7
105 <211> LENGTH: 8
106 <212> TYPE: PRT
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: peptide in m13 coliphage
112 <400> SEQUENCE: 7
      Tyr Thr Arg Asp Phe Glu Ala Gly
114
115
117 <210> SEQ ID NO: 8
118 <211> LENGTH: 8
119 <212> TYPE: PRT
120 <213> ORGANISM: Artificial Sequence
122 <220> FEATURE:
123 <223> OTHER INFORMATION: peptide in ml3 coliphage
125 <400> SEQUENCE: 8
127
     Ser Ser Ala Ala Gly Pro Arg Gly
128
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130 <210> SEQ ID NO: 9
131 <211> LENGTH: 8
132 <212> TYPE: PRT
133 <213> ORGANISM: Artificial Sequence
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Input Set : A:\46004seq.txt

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    136 <223> OTHER INFORMATION: peptide in m13 coliphage
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    145 <212> TYPE: PRT
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    152 <221> NAME/KEY: MISC_FEATURE
    153 <222> LOCATION: (7)..(7)
    154 <223> OTHER INFORMATION: unknown
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    159
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    163 <212> TYPE: PRT
    164 <213> ORGANISM: Artificial Sequence
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    170 <221> NAME/KEY: MISC_FEATURE
    171 <222> LOCATION: (3)..(3)
    172 <223> OTHER INFORMATION: unknown
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WALL $ 176
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    177
           1
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    181 <212> TYPE: PRT
    182 <213> ORGANISM: Artificial Sequence
    184 <220> FEATURE:
    185 <223> OTHER INFORMATION: peptide in ml3 coliphage
    187 <400> SEQUENCE: 12
          Phe Asn Thr Ser Thr Arg Thr Gly
    189
            1
    192 <210> SEQ ID NO: 13
    193 <211> LENGTH: 8
    194 <212> TYPE: PRT
    195 <213> ORGANISM: Artificial Sequence
    197 <220> FEATURE:
    198 <223> OTHER INFORMATION: peptide in ml3 coliphage
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     202
     203
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Input Set : A:\46004seq.txt

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     210 <220> FEATURE:
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     213 <400> SEQUENCE: 14
     215
          Asp Tyr Ser Phe Pro Pro Leu Gly
     216
           1
     218 <210> SEQ ID NO: 15
     219 <211> LENGTH: 8
     220 <212> TYPE: PRT
     221 <213> ORGANISM: Artificial Sequence
     223 <220> FEATURE:
     224 <223> OTHER INFORMATION: peptide in ml3 coliphage
     226 <400> SEQUENCE: 15
    228
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     229
     231 <210> SEQ ID NO: 16
     232 <211> LENGTH: 8
    233 <212> TYPE: PRT
     234 <213> ORGANISM: Artificial Sequence
     236 <220> FEATURE:
     237 <223> OTHER INFORMATION: peptide in m13 coliphage
    239 <220> FEATURE:
    240 <221> NAME/KEY: MISC FEATURE
    241 <222> LOCATION: (2)..(2)
     242 <223> OTHER INFORMATION: unknown
     244 <220> FEATURE:
    245 <221> NAME/KEY: MISC_FEATURE
     246 <222> LOCATION: (6)..(6)
     247 <223> OTHER INFORMATION: unknown
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    252
            1
    254 <210> SEQ ID NO: 17
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    256 <212> TYPE: PRT
    257 <213> ORGANISM: Artificial Sequence
    259 <220> FEATURE:
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    265
           1
    267 <210> SEQ ID NO: 18
    268 <211> LENGTH: 8
    269 <212> TYPE: PRT
    270 <213> ORGANISM: Artificial Sequence
    272 <220> FEATURE:
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Input Set : A:\46004seq.txt

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278
280 <210> SEQ ID NO: 19
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282 <212> TYPE: PRT
283 <213> ORGANISM: Artificial Sequence
285 <220> FEATURE:
286 <223> OTHER INFORMATION: peptide in ml3 coliphage
288 <400> SEQUENCE: 19
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291
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293 <210> SEQ ID NO: 20
294 <211> LENGTH: 8
295 <212> TYPE: PRT
296 <213> ORGANISM: Artificial Sequence
298 <220> FEATURE:
299 <223> OTHER INFORMATION: peptide in m13 coliphage
301 <400> SEQUENCE: 20
303
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304
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306 <210> SEQ ID NO: 21
307 <211> LENGTH: 8
308 <212> TYPE: PRT
309 <213> ORGANISM: Artificial Sequence
311 <220> FEATURE:
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314 <400> SEQUENCE: 21
316
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317
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319 <210> SEQ ID NO: 22
320 <211> LENGTH: 8
321 <212> TYPE: PRT
322 <213> ORGANISM: Artificial Sequence
324 <220> FEATURE:
325 <223> OTHER INFORMATION: peptide in m13 coliphage
327 <400> SEQUENCE: 22
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330
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334 <212> TYPE: PRT
335 <213> ORGANISM: Artificial Sequence
337 <220> FEATURE:
338 <223> OTHER INFORMATION: peptide in m13 coliphage
340 <400> SEQUENCE: 23
342 Thr Thr Pro Pro Asn Phe Ala Gly
343
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Input Set : A:\46004seq.txt

Output Set: N:\CRF4\10072002\J053520.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:10; Xaa Pos. 7
Seq#:11; Xaa Pos. 3
Seq#:16; Xaa Pos. 2,6
Seq#:47; N Pos. 19
Seq#:48; N Pos. 8
Seq#:53; N Pos. 6,16,18
Seq#:100; Xaa Pos. 5
Seq#:133; N Pos. 14
Seq#:251; Xaa Pos. 7
Seq#:252; Xaa Pos. 3
Seq#:257; Xaa Pos. 2,6
Seq#:304; Xaa Pos. 5